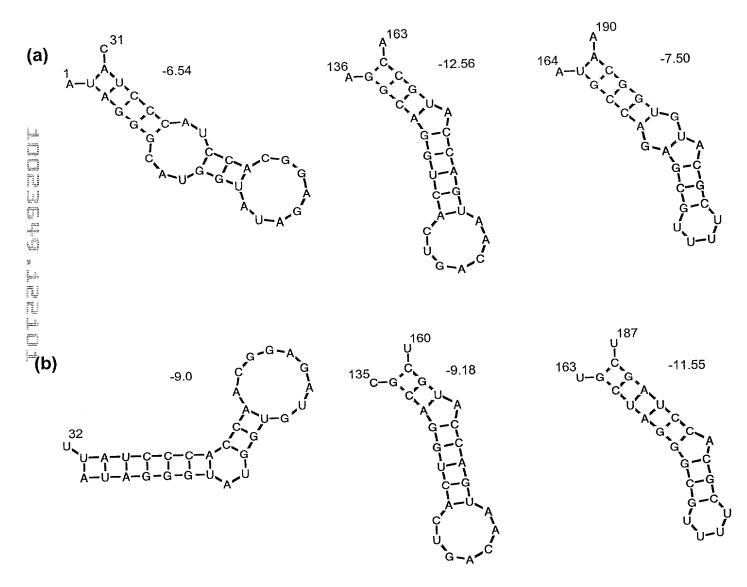


Figure 2 Multiple sequence alignment

(a)	1 1 1 1	A T A	 A G	 G	 G I	- A	<u></u>	 : G	<u></u>	- ; T	A					- <u>-</u>	_ C	_ Cl	A (	2 C	C	T.	A C	: <u>A</u>	A T	A C	T.	A P	A I C	CBA87 71V-1658 EEE VEE
	41 17 39 37	GATA	r c r c	C	A A	TA	r A r A	' G	G A	AG	A (	G A	A	T	T (	A A	C	G G	T I	ľG ľG	A	C '	r 1 r 1	A A	G.	AT	G	C I	G A G A A G A	CBA87 71V-1658 EEE VEE
	81. 57 79 77	CA	G C	C	C G	T Jr[	<u>А</u> Т	G	T C	A A	A (	G I G I	' C	A	cl:	rG	c	Α	A A	AG	A	T	G (	T	Т	T C	C	<b>A</b> (	A A A T A G	CBA87 71V-1658 EEE VEE
of control		TTT	ľ G	Α	$G_{\underline{P}}$	T	A	A	ΑG	C.	A	A C	: G	C.	A (	G G	Т	С	A (	CT	G	A	C, I	A	Т	G A	C	C I		71V-1658 EEE VEE
	159	C C T	A A	$\mathbf{T}$	G C		A G	;[G	GC	G	T	r 1	T	С	G (	: A	C	C	ΤZ	A G	c	T[	A C	T	A	A_G	C	T_C	CAT	71V-1658 EEE VEE
	97 177 199 197	TGI TGI	A G	G	G	JG	A A	4 G	TIG	ΙG	A	r A	C	A	G	A C	C	A	G	G T	G	Α	T (	cc	T	GG	A	TI	AT ATT ATT	71V-1658 EEE VEE
(b)	C '	I C G C T G T C G	A C	ГА ГА Г <u>А</u>	T	G G A G	G G	C I G C A I	TT	C (T) (G) (G)	G C C G	C G T	C (	G T G T	A	G (G	G C G T A A	T C C T T	C C	A A[ A	A G	50 50		V-1 EE-						

- a. The 5' terminus of WEE CBA87 (1-97), WEE 71V-1658 (25-240), EEE (1-238) and VEE (1-236) via Clustal module of DNAStar. Areas where sequences differ are boxed.
- b. Hypervariable region identified in nsP1. Alignment of WEE 71V-1658 (1420-1449), WEE 1654 (65-94) and EEE (1415-1444) is shown.

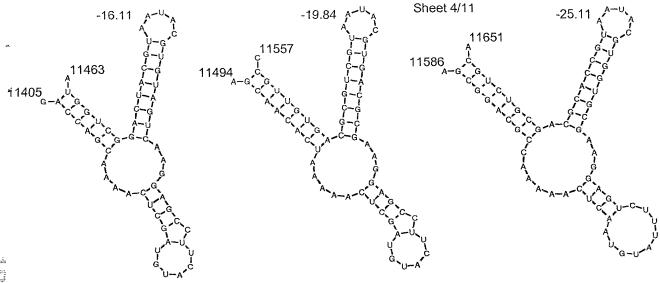
Stem loop structures in the 5' NTR Figure 3



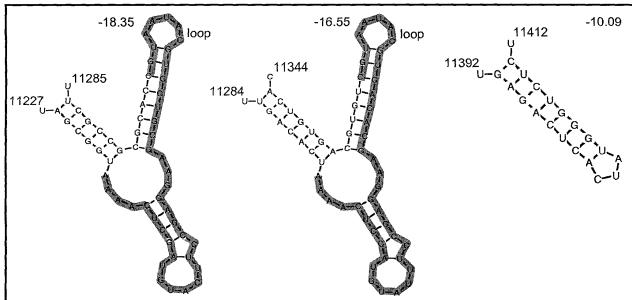
Hairpin structures were identified using the RNA folding program of the Genequest module (DNASTAR).

- a. Structures for WEE (CBA87/71V-1658) sequence (1-192).
- b. Structures for EEE (1-192).

Minimal free energy values are shown for the different structures.



Double stem loop structures in SIN.



(b) Double stem loop structures in 3' NTR of WEE. Residues in the SIN-like 40 nt repeat are shaded.

Figure 4 Stem loop structures in the 3' NTR

Figure 5 Phylogenetic relationship of the WEE nonstructural region compared to other alphaviruses

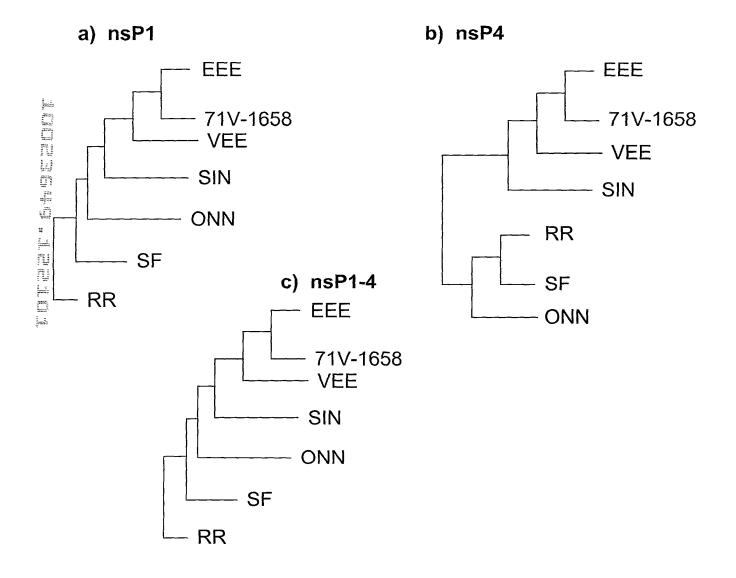
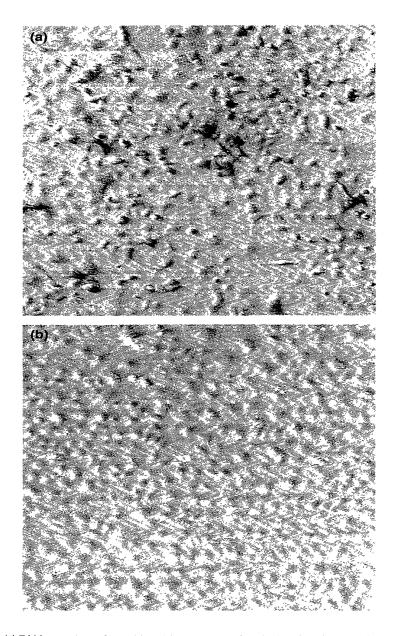
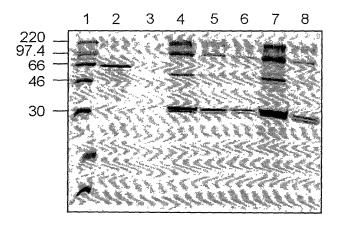


Figure 6 Expression of WEE structural genes in cell culture



One  $\mu g$  of plasmid DNA was transfected into Vero cells. After 31 hrs incubation, the cells were histochemically stained using a monoclonal antibody to WEE (11D2). a. pCXH-3; b. pCl (control plasmid).

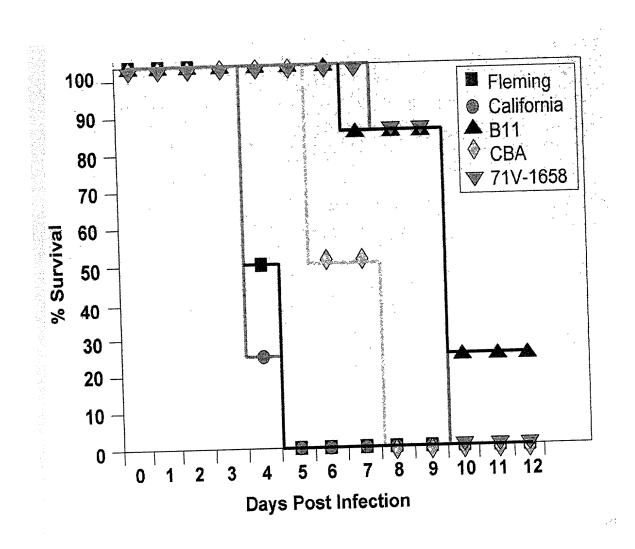
Figure 7 In vitro transcription and translation of WEE expression vectors



Qiagen purified vectors containing the WEE 26S insert were expressed *in vitro* using the TNT system and [ $^{35}$ S]-methionine labelling. Three  $\mu$ L aliquots of each samples were run by SDS-PAGE on a 12% gel.

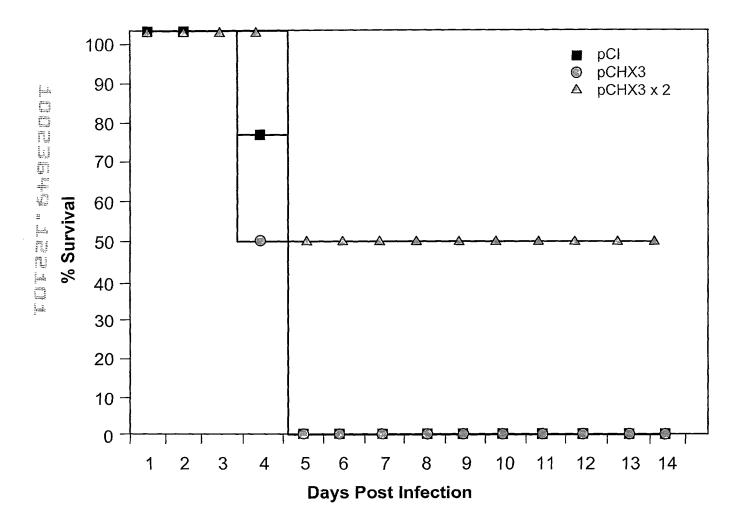
Lane: 1) Rainbow <sup>14</sup>C-labelled marker; 2) Luciferase translation control; 3) pVAX; 4) pVHX-6; 5)pCXH-3; 6) pcDWXH-7; 7) pcDWHX-45; 8) pXTR2-4.

Figure 8 WEE mouse infectivity model



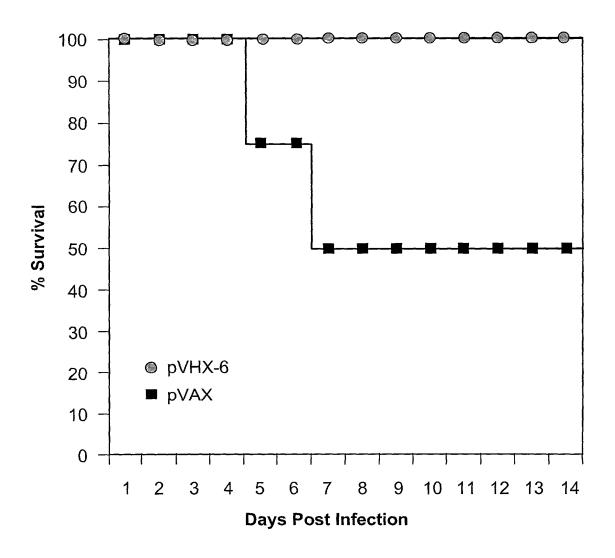
Groups of 4 mice were inoculated intranasally with 50  $\,\mu$ L of virus (approximately 10<sup>4</sup> PFU). The mice were monitored for 12 days, and the % survival graphed.

Figure 9 Protection using ballistic delivery of pCXH-3



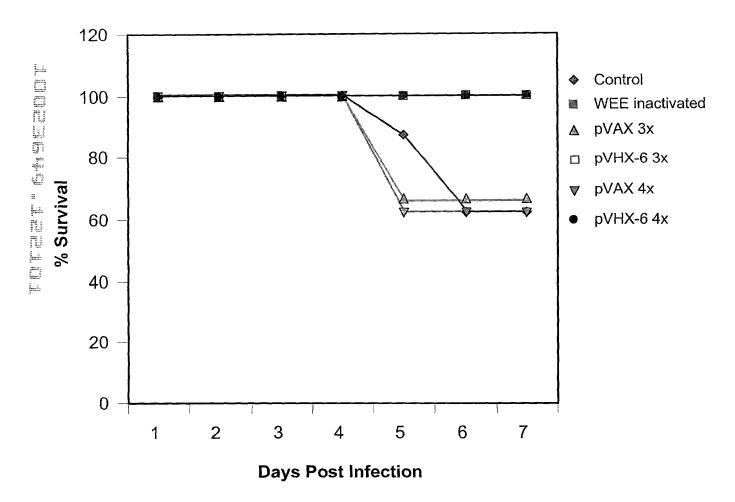
Groups of 4 mice were immunized with one or two doses (2 x 1.25  $\,\mu$ g) of either pCl or pCXH-3. The interval between boosters (2 doses) or challenge was 3 weeks. The mice were challenged intranasally with 50  $\,\mu$ L of WEE Fleming (1.25 x 10<sup>4</sup> PFU). The mice were monitored for 12 days, and the % survival graphed.

Figure 10 Protection using ballistic delivery of pVHX-6



Groups of 4 mice were immunized with four doses (2 x 1.25  $\,\mu g)$  of pVAX or pVXH-6 . The interval between boosters or challenge was 2 weeks. The mice were challenged intranasally with 50  $\,\mu L$  of WEE Fleming (1.25 x 10^4 PFU). The mice were monitored for 14 days, and the % survival graphed.

Figure 11 Protection using ballistic delivery of pVHX-6



Groups of 5-8 mice were immunized with three or four doses (2 x 1.25  $\,\mu g)$  of pVAX or pVXH-6 . The interval between boosters or challenge was 2 weeks. The mice were challenged intranasally with 50  $\,\mu L$  of WEE Fleming (1.7 x 104 PFU). Untreated control and WEE inactivated control (3 doses) groups were also included. The mice were monitored for 14 days, and the % survival graphed.